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PEPTIDESTRUCTURE of: human_prot.pep ck: 2321, 1 to 458

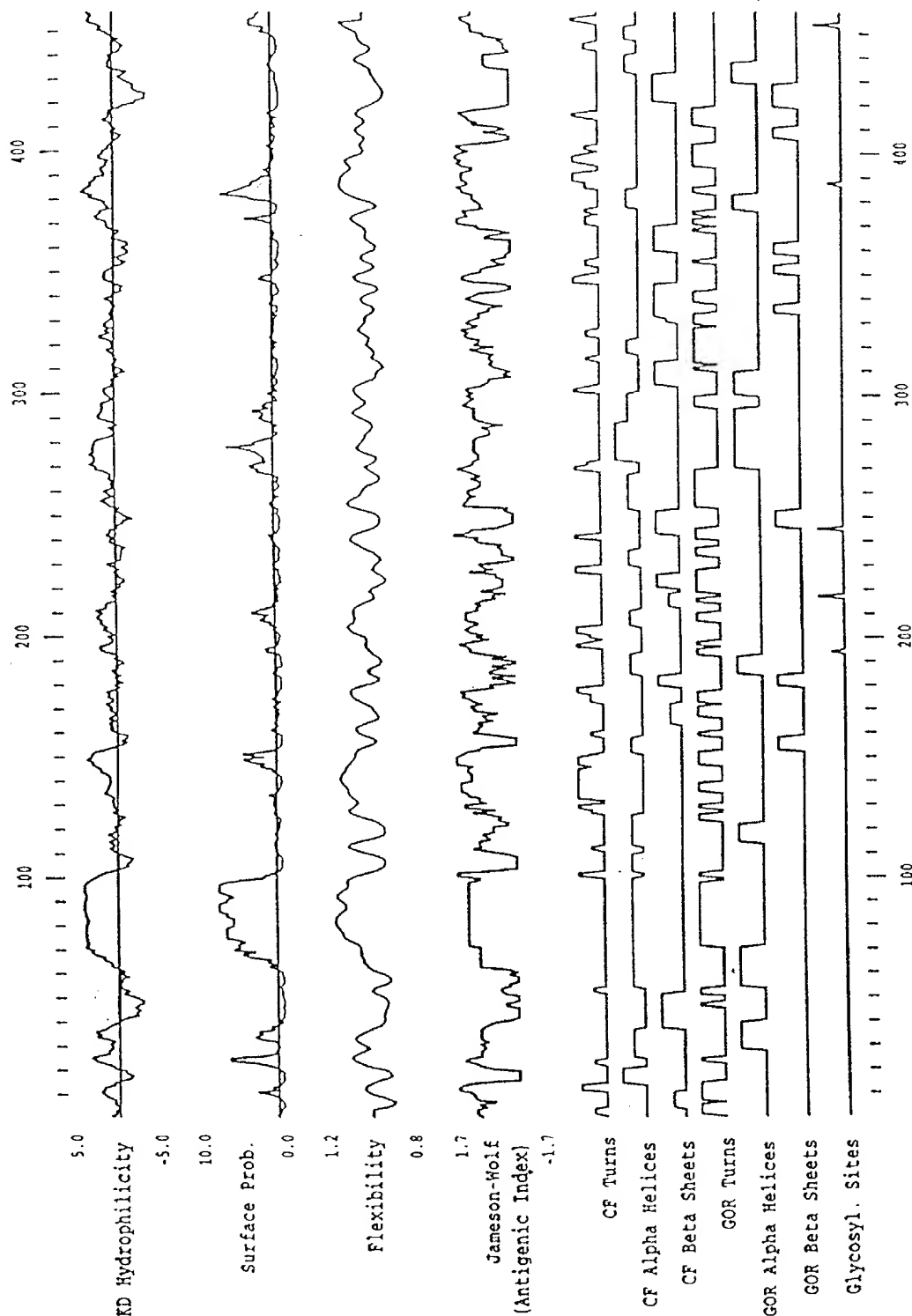
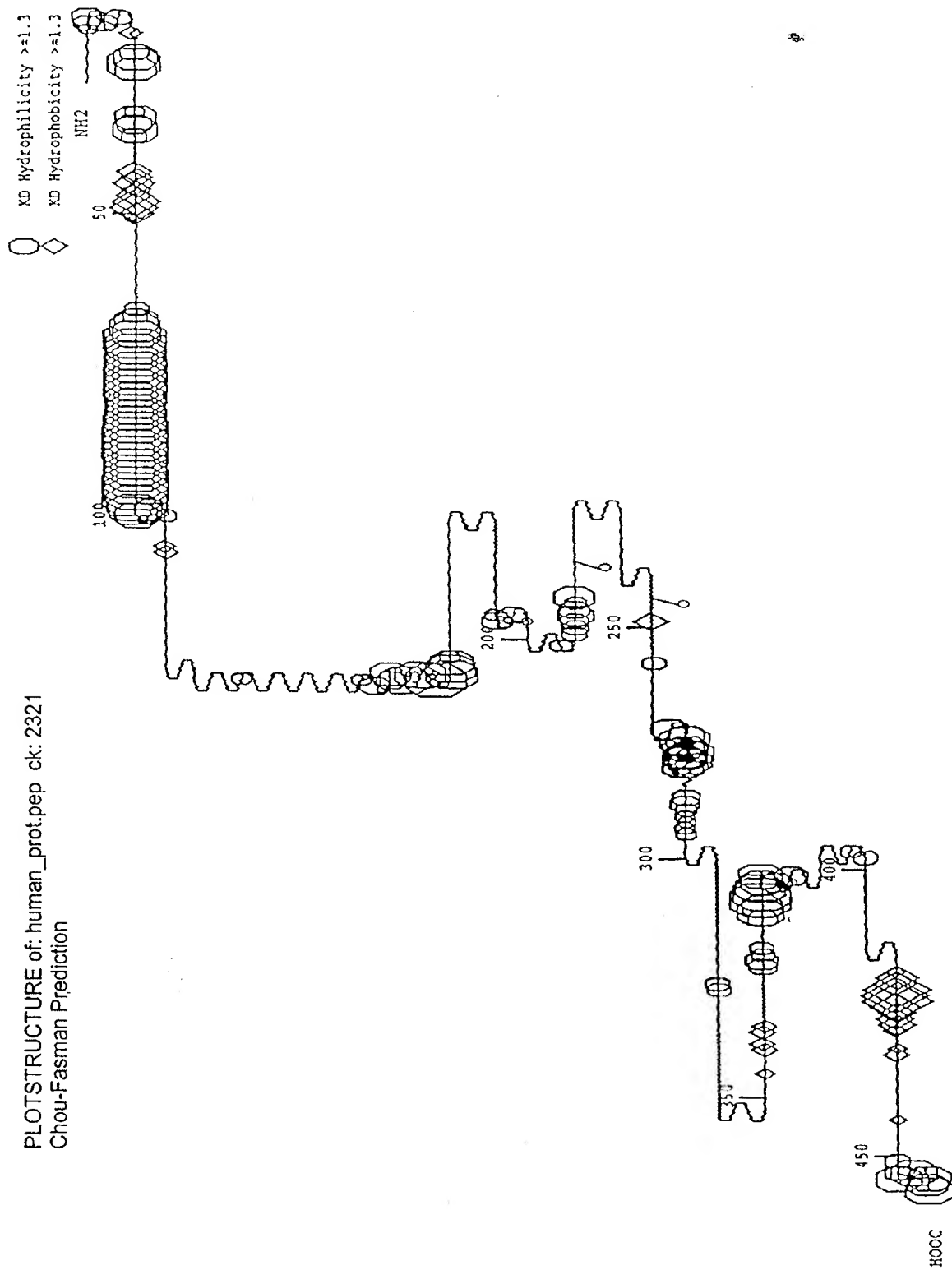


FIGURE 1

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PLOTSTRUCTURE of: human_prot.pep ck: 2321
Chou-Fasman Prediction

FIGURE 2

FO9060 90520860

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KD Hydrophilicity >=1.3
KD Hydrophobicity >=1.3



PLOTSTRUCTURE of: human_prot.pep ck: 2321
Garnier-Osguthorpe-Robson Prediction

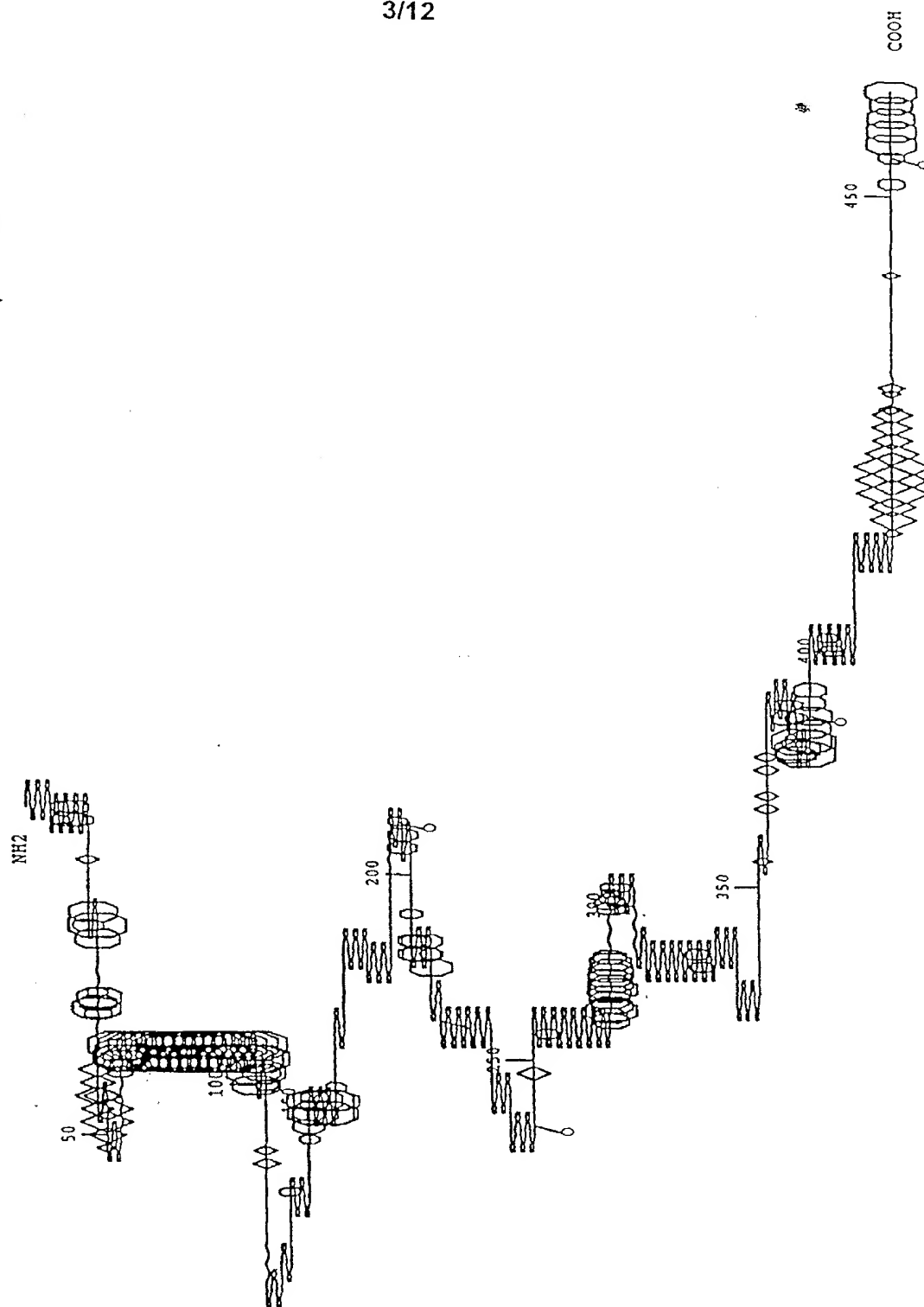


FIGURE 3

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PEPTIDESTRUCTURE of: g713estmouse.0.12_tr13.pep ck: 1880, 1 to 467

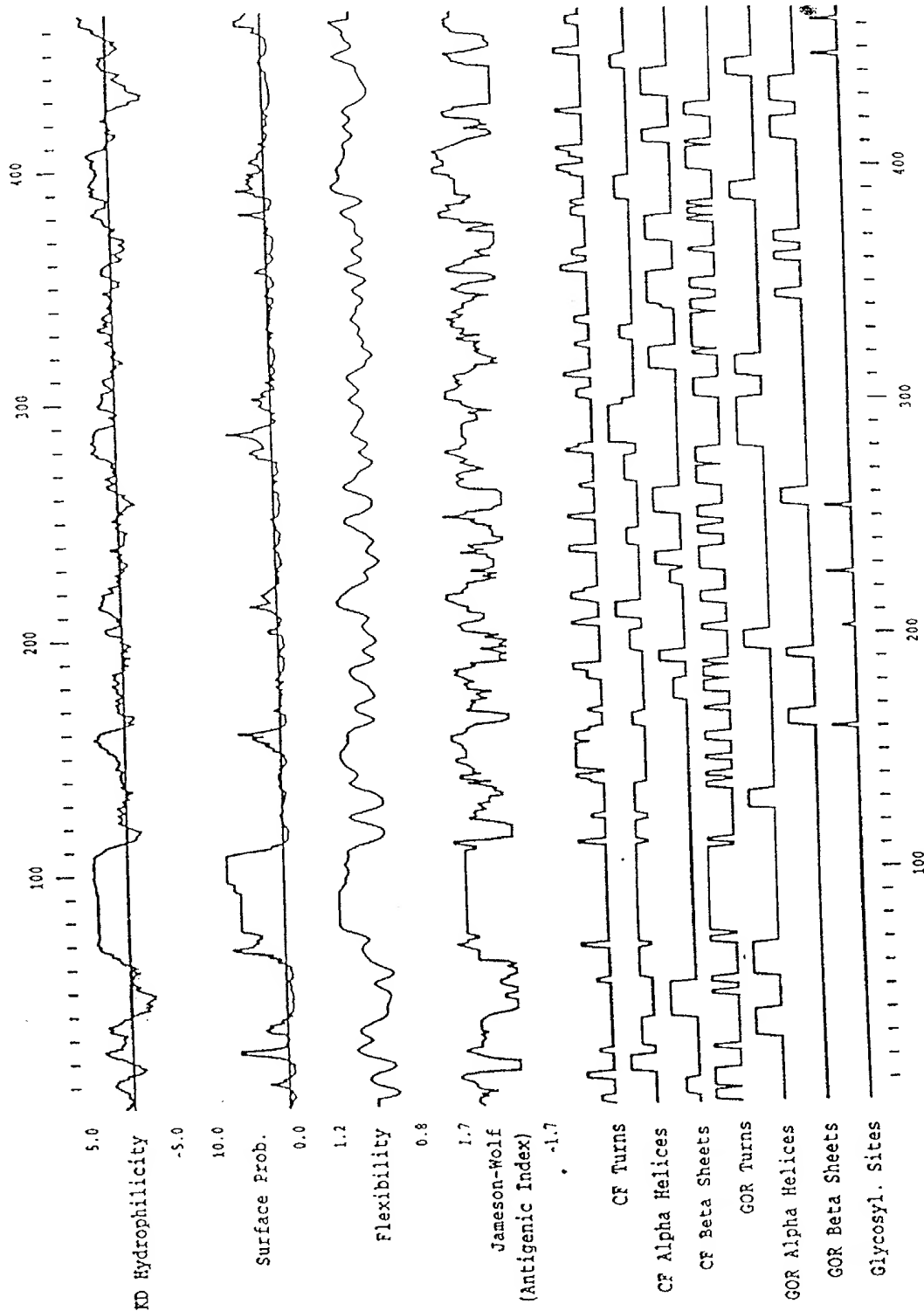


FIGURE 4

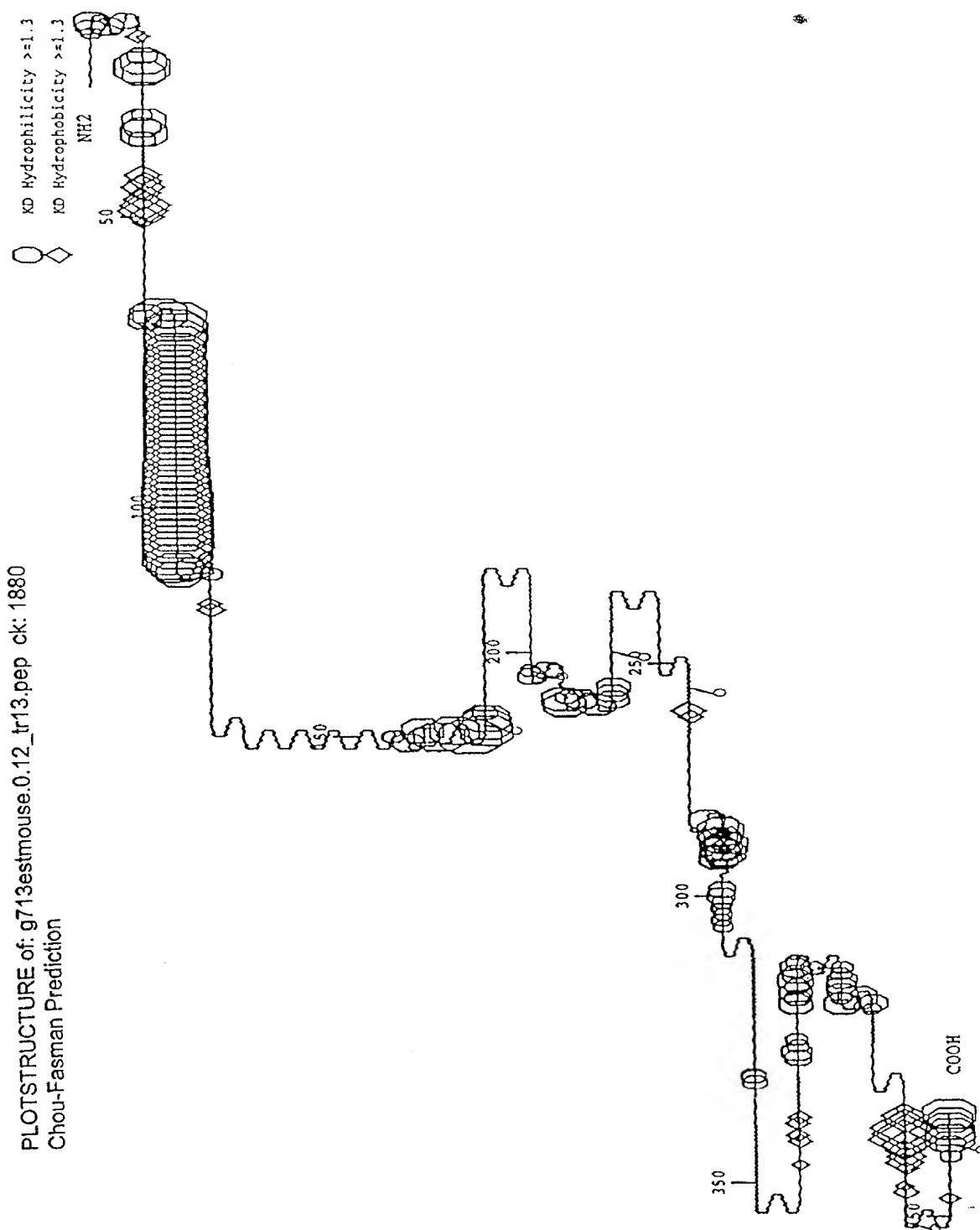


FIGURE 5

T09080" 909/0860

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XD Hydrophilicity >=1.3
XD Hydrophobicity >=1.3



PLOTSTRUCTURE of: g713estmouse.0.12_tr13.pep ck: 1880
Garnier-Osguthorpe-Robson Prediction

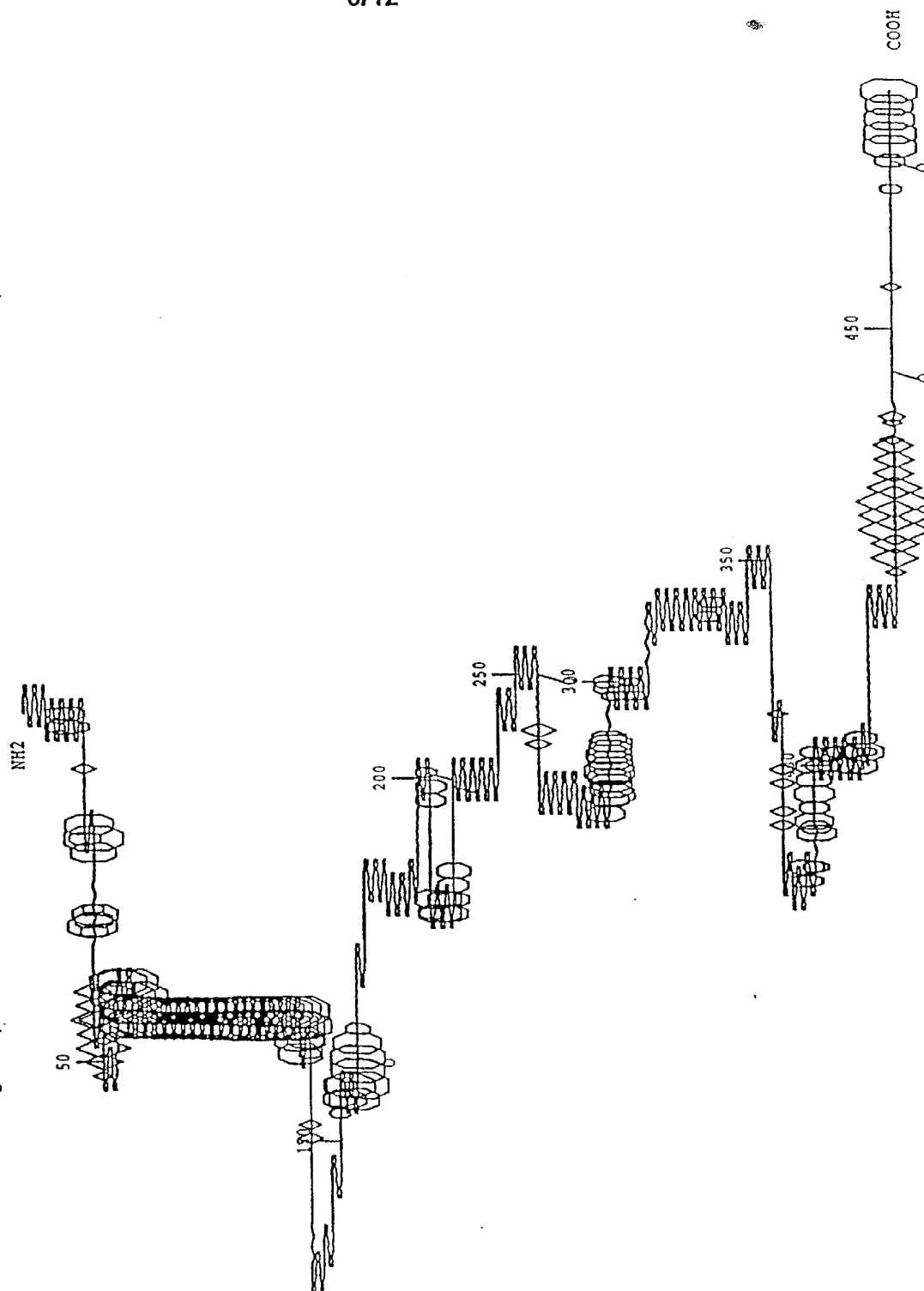


FIGURE 6

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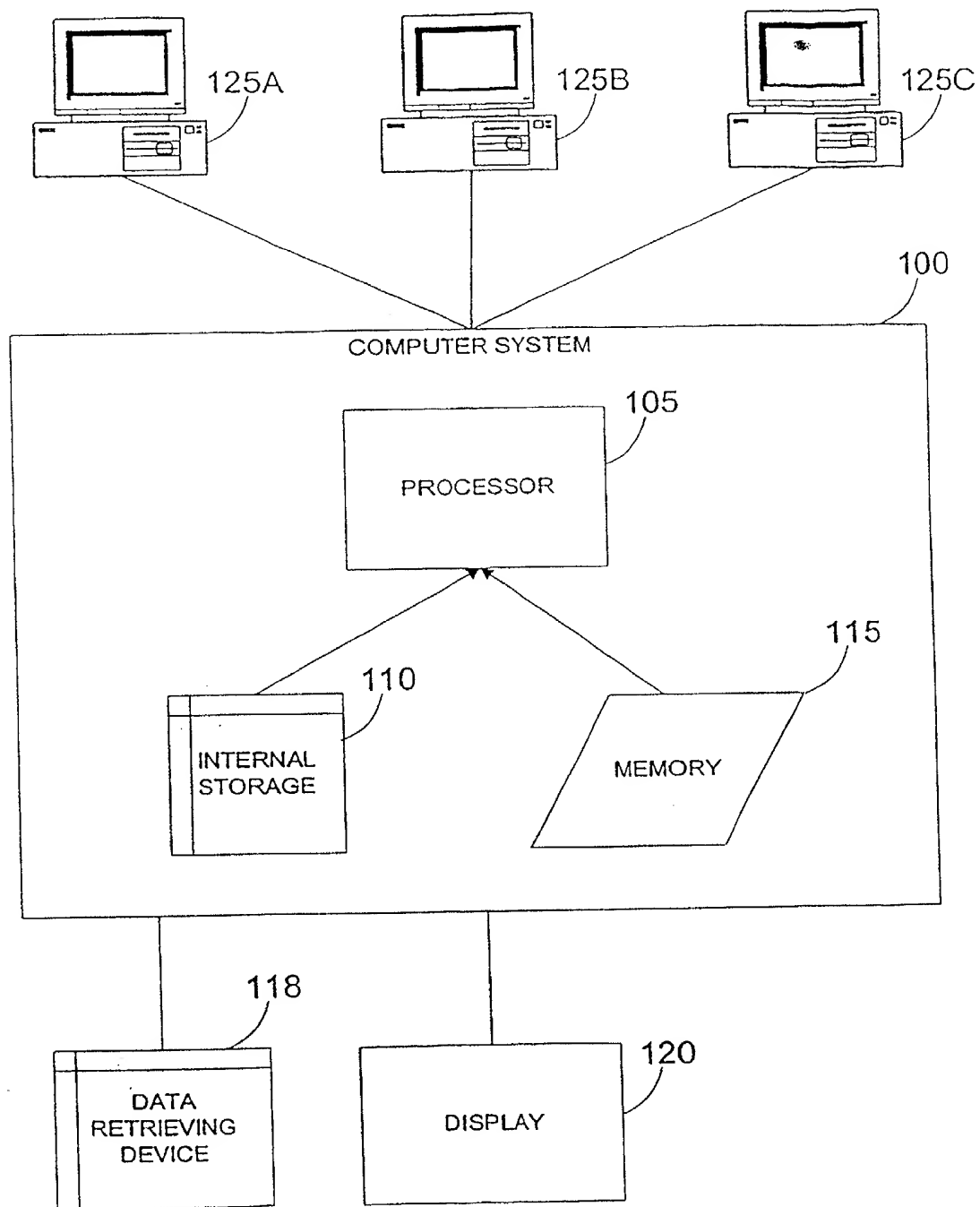


FIGURE 7

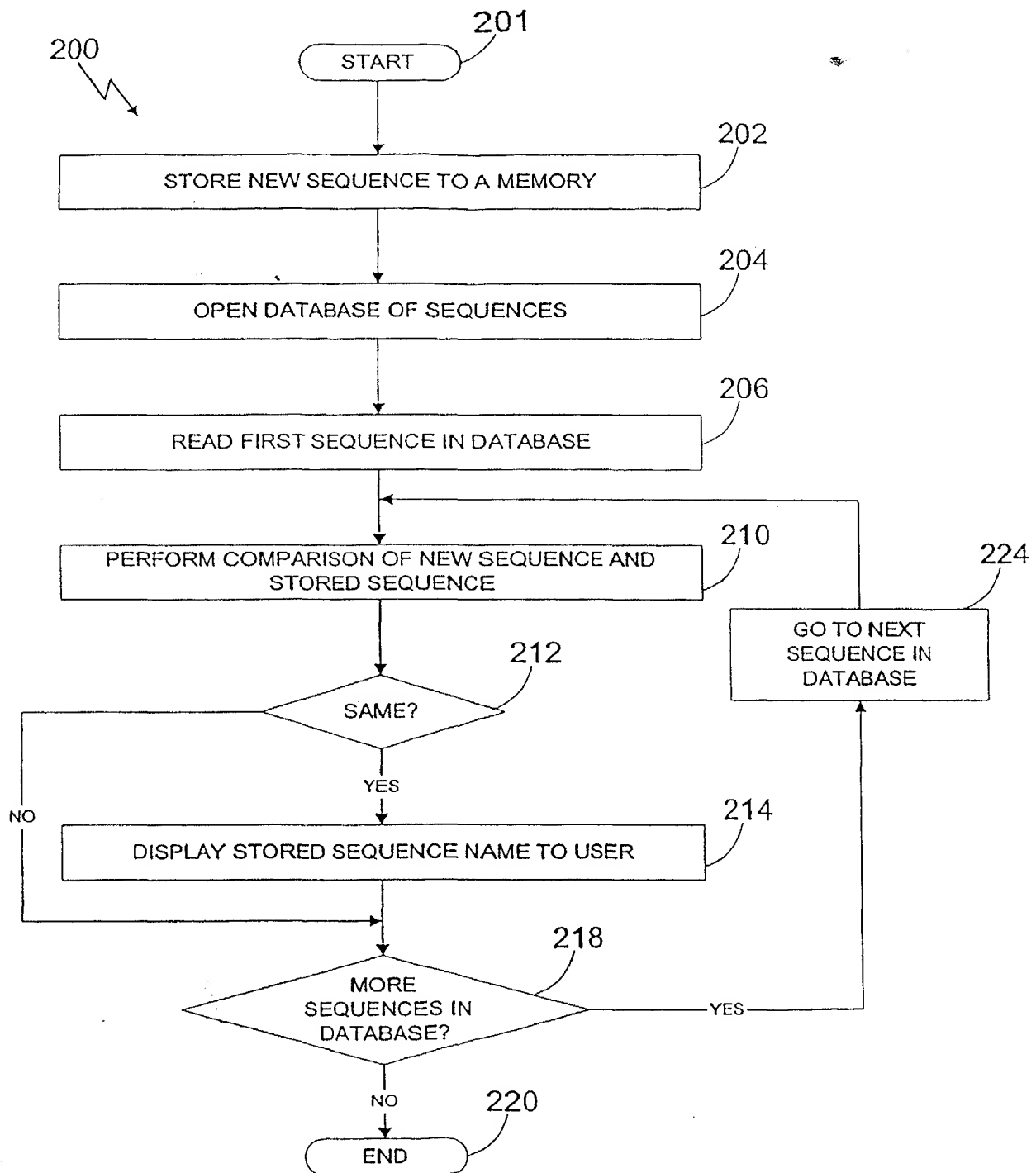


FIGURE 8

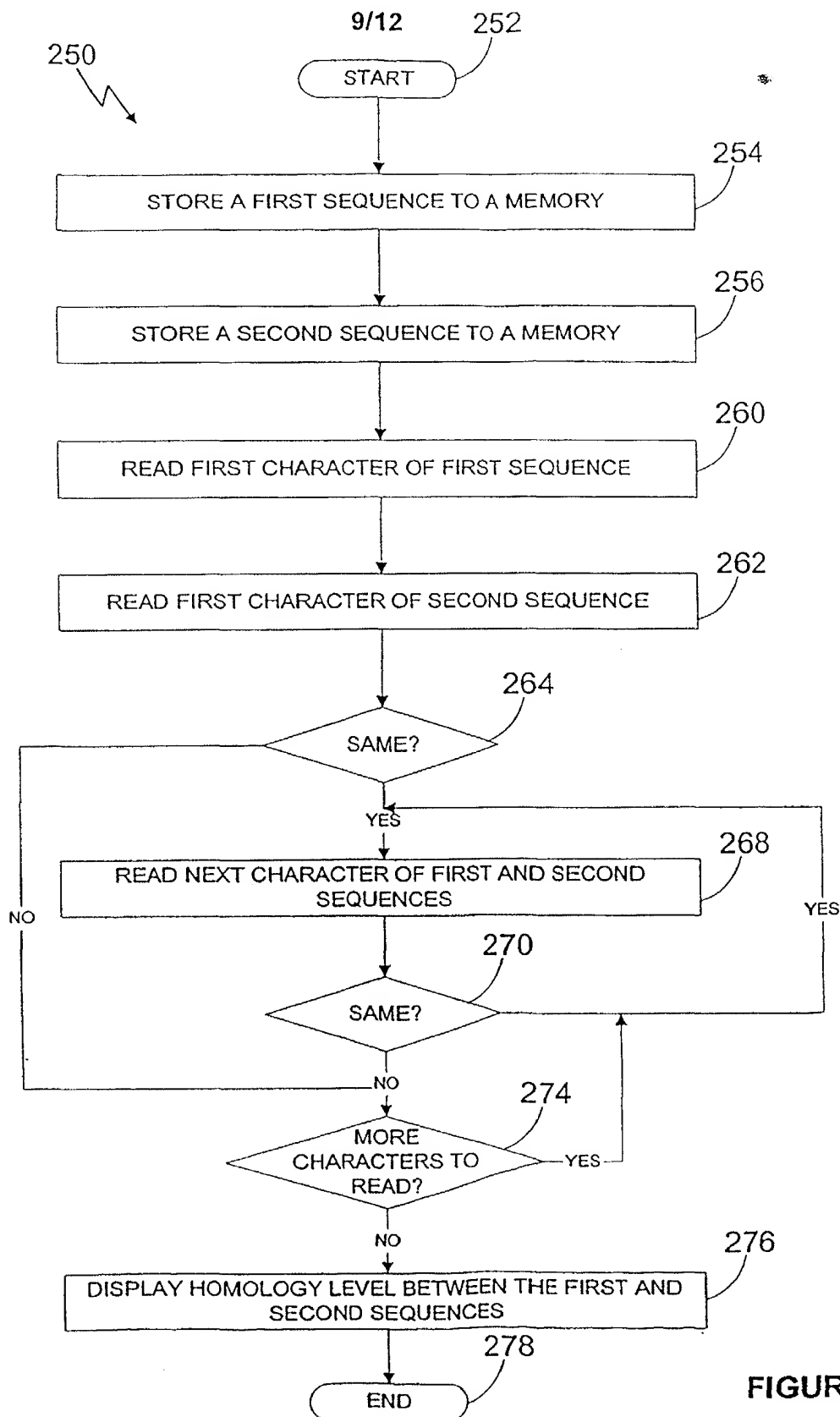


FIGURE 9

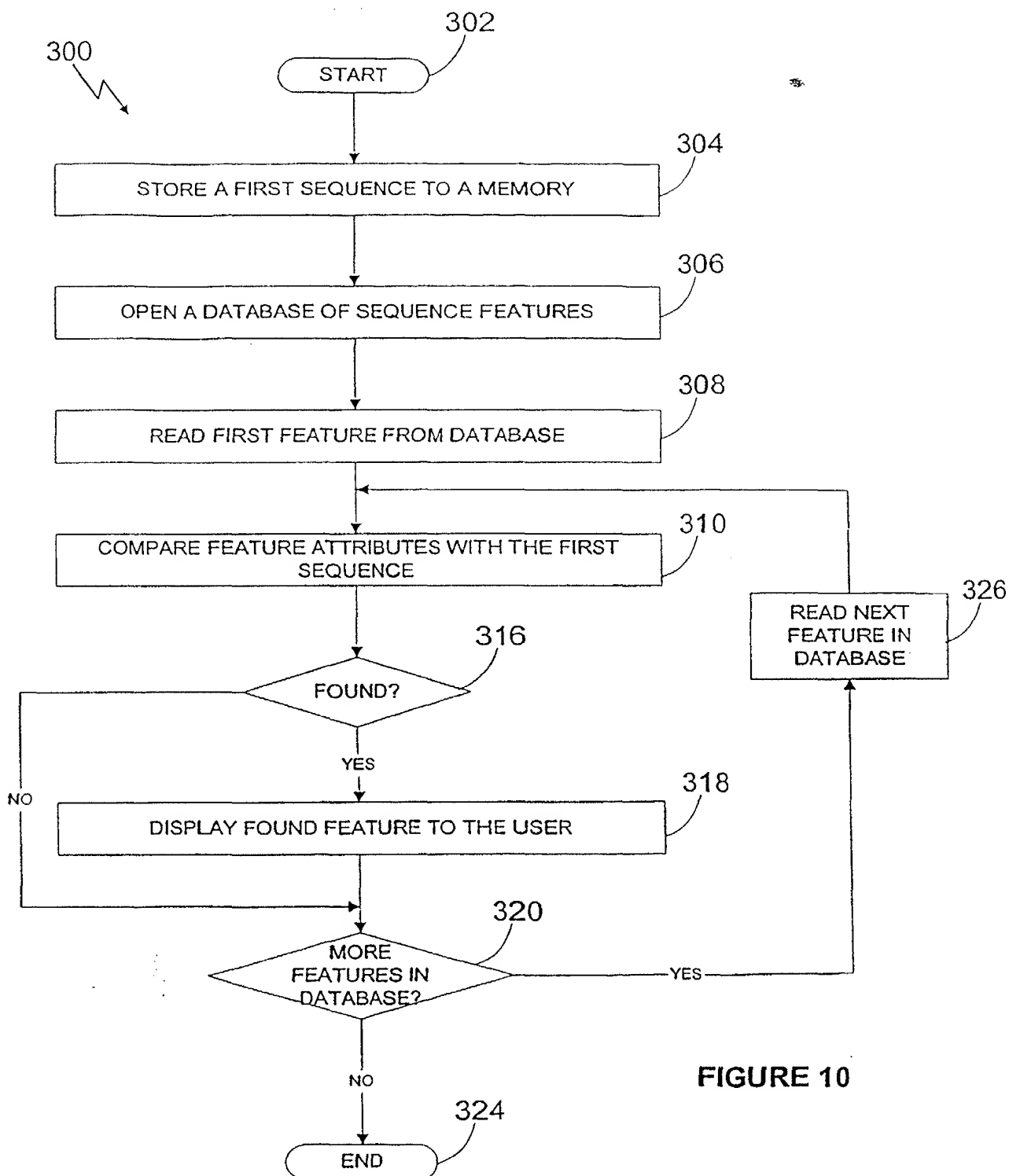


FIGURE 10

FIGURE 11

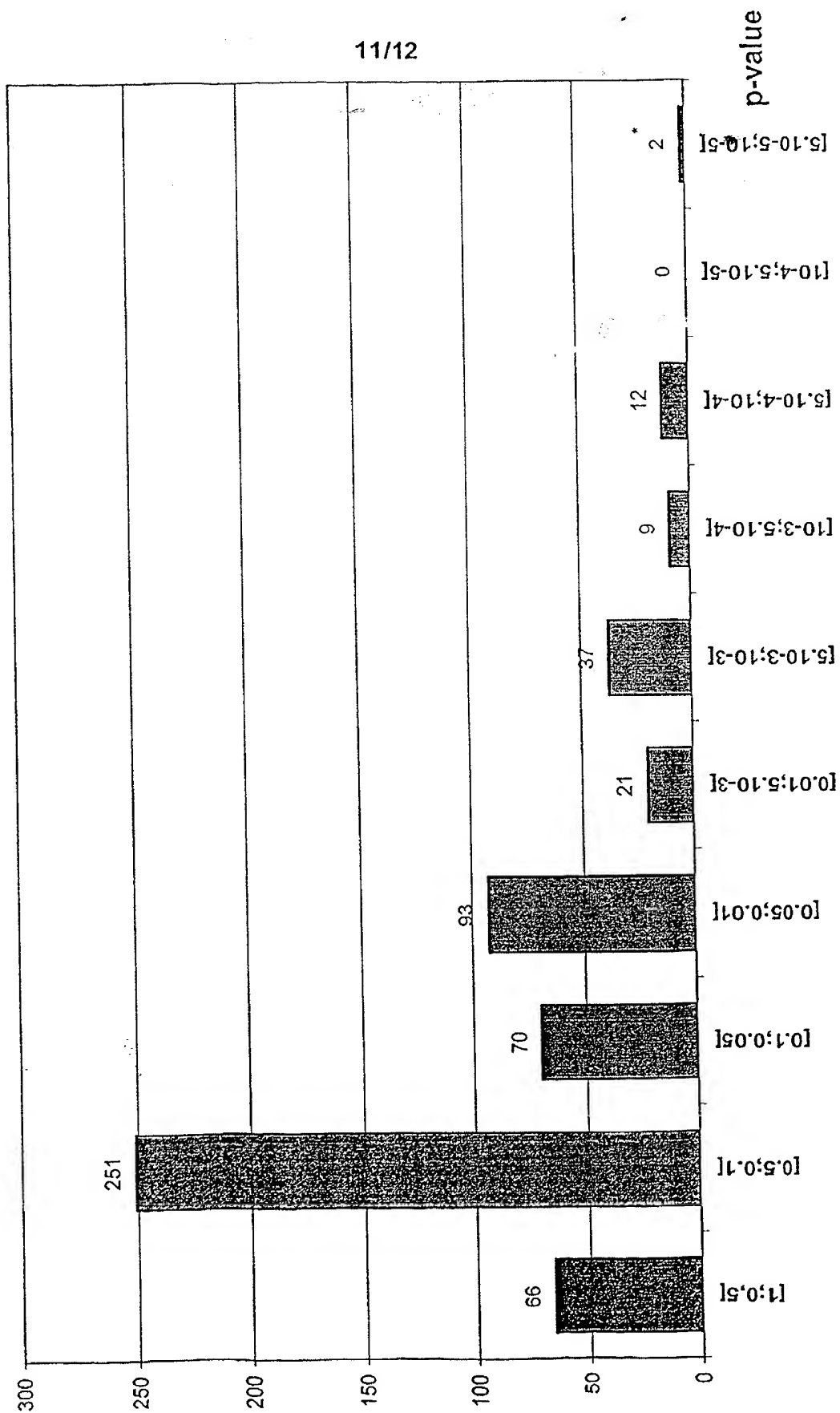


FIGURE 12

